

RAW SEQUENCE LISTING

**The Biotechnology Systems Branch of the Scientific and Technical
Information Center (STIC) no errors detected.**

Application Serial Number: 10/578,158
Source: IFWP
Date Processed by STIC: 05/16/2006

ENTERED



IFWP

RAW SEQUENCE LISTING

DATE: 05/16/2006

PATENT APPLICATION: US/10/578,158

TIME: 09:46:53

Input Set : A:\E5106-00003 Motoneuronotrophic Factor Gene Sequences.txt

Output Set: N:\CRF4\05162006\J578158.raw

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3 <110> APPLICANT: XUE, Bob, Baoguo
5 <120> TITLE OF INVENTION: Motoneuronotrophic Factor Gene Sequences
7 <130> FILE REFERENCE: E5106.00003
C--> 9 <140> CURRENT APPLICATION NUMBER: US/10/578,158
C--> 9 <141> CURRENT FILING DATE: 2006-05-04
9 <150> PRIOR APPLICATION NUMBER: 60/518,581
10 <151> PRIOR FILING DATE: 2003-11-07
12 <160> NUMBER OF SEQ ID NOS: 32
14 <170> SOFTWARE: PatentIn version 3.3
16 <210> SEQ ID NO: 1
17 <211> LENGTH: 1859
18 <212> TYPE: DNA
19 <213> ORGANISM: Homo sapiens
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26 acaactgttt tctgtgccc aagagagttg caagagcacc ttgaaaacac aagcaaacat 180
28 atcatatcaa cttccaggca tctaattaat tgtgcaccaa aatcgttacc cttccaatct 240
30 tgcaacagaa gagcaaaagc tgcttagacc ttactcaaag gctccaaagc agaaacaagt 300
32 tttgcttctt gcagcaatgt gtaacagagg aaacagatgt cctggaagat cccatggtct 360
34 gtttcaggtg agatggaacc aatgctgcac acgaagattc atctcaaag aaatgcacag 420
36 gaaaagccaa ttactttatg tgaataaaag aataaatccc taaagcagtg gttctcaacc 480
38 agagtgtac caccacacc ccagagggca tttgggaatt cttggggaca ttttggggtg 540
40 acacactgaa ctgctggatg ctatcagcat ttagtaggta tgctcgatgt cttgcagaag 600
42 gacatgatgg tcctacacag taaggaatgg attacctaca atattaatag cagcctccca 660
44 tacacacttt tgacaccctt ccctaaagga ttaatatgct ccaaccttcc tgtccccaca 720
46 gttcagtggt tctccctacc ctcaccatga tcggatgaaa aaaaataagg tttcacagct 780
48 taagagttaa attctggaat ccaactacaa gctcataact gtagcatgga acctggtagt 840
50 agcataataa ataaattttt agtaagagge ttaagaaatt ttagcaaaaa aagcactccc 900
52 tttcttcttc cctacatata tcatatgttt ttcaacacaa aaaattctgt gatttttagag 960
54 aaactttctt cagtactttt aagttcaaaa ccagatgctc attacagttc ttttaaacac 1020
56 caaactagtc atctcaaaaa tatggctaac tctctggact aaattccata ggaaaaatta 1080
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78 ttgctggcct gtcaccagaa gttcataaaa tgaattcatc cagattctct gaatcctcct 1740

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80 tttccccagt catgtgccaa cagccaggca ataatgcccc accgtgacag gctccctgtt 1800
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86 <211> LENGTH: 4359
87 <212> TYPE: DNA
88 <213> ORGANISM: Homo sapiens
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95 agctaacaat aaaagaaaat taaatcactt gagaacccta cagtggcagc aaaatagagc 180
97 ccaagagata cctcctagca tgttcaaaga cttattaaaa gccacgttgg gccaggcgtg 240
99 gtggcttatg cctataatcc caataccttg ggaggttgag gtgggaggat tgcttaaagc 300
101 cggagtttg aggccagcct aggcaacata gtgagacctc atctctacaa aagattgtat 360
103 aaaatttagc caggtgtagt ggtgcgcacc tgagtccag ctactctgga agctgagatg 420
105 ggaagacggc ttgagccaa gagtttgagg ctacagtga ccatgattgc accatggcac 480
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109 cacaatcacc ttgtatcagt ctcagttgat ggcttgatga caataaagaa cagtttagac 600
111 tttctgaaag tgcatacata tatatacaca tatgactaag aagggtgaagt gaagccctca 660
113 atggccagcc cttttatagt agaacctaga aaacagcaac ttctttcaac atgggtggatg 720
115 gtgcatgttt tgtacccac atttagaaat ccatactctg agacaaacag aactaccgaa 780
117 agagaatgac caggcctcca gaatagcacc gaggcagcca tcaggagtta tagcaaaact 840
119 ttttaacgtg gaaagggagt cactctgctt ctataaagca gcccttatca gcggcattat 900
121 gagtaaaagc agcccactaa tgcaggccag actttaggac acagaaccag attccatttg 960
123 ggggctacag ctagcacagc tgtaaacctc tctccacca tcaaattctg agaccattag 1020
125 gcctggcacg gtggctcatg cctgtggtcc cagcactttg ggaagccaag ctgggaggat 1080
127 caactgagcc caggagtctg agactgcagt gagctatgat ctcgccactg cactccagcc 1140
129 tgggtgacag aataagaccc tgtctcaaaa gaaaaaaaaa aaaaatctga ggcgattgaa 1200
131 ggcagacaac tataagagag tgccccata ggaactccac cactctccat ttagaatgtc 1260
133 ctttgacttt tgtgcacatt atctcatgag accctcacta tgactctgag aggtaagcag 1320
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137 tgcccaaggt cacaaatcat gcagctaate agctgcctaa ccaggagatg agcctagcac 1440
139 tctgactcgt aggccagcgc aatctccagg acaccatgcg gctttgtgat gttttcagat 1500
141 aagaggcgga agaggggtact ggttaagagc ccagggtctg gcaccagcca acctgcattc 1560
143 caatcccatc tccaccacct actaaccacc tgatcttggtg tgagtccctt aacctcccca 1620
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151 cggagcttag actttaggca aaccatacac agaaaaaagt aaacagataa aatacttaca 1860
153 aagtgtgtta agaattgatta aggtgcaggg cgcagtggct cagcctgta atcctatcac 1920
155 tttgggaggg cggttcaggc agatcacctg aggtcgggag tttgagacca gcctgataaa 1980
157 catggagaaa tcccgctctc acaaaaaaca aaattagccc caagcttatt gggcattggtg 2040
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161 gaggcgagga ctgcagttag cagaggtcgc accactgcac tccagcctgg gcaacaagag 2160
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165 accggagggg ccaattttta agagtggctc ctaagcccta agggatcaga aagagaaatt 2280
167 cgtgtgaaga acggagagaa aagcactcta ggattagggg atgggtatacg tgggggtcac 2340
169 agcacaggga agtcaggaac ctagagaaga caaatgtgac tacagcgctt cctctcccg 2400
171 gatttgcgga gcaaggctgg agctcgtgag ggtgaccaca actattttta aaacgacaaa 2460
173 gataaggctc aagagagggg atatgtcttg cccaaggcca cagctgctca gaggagtagc 2520

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175 tgggaccagg gcccttcagt attctacctc cagaccacag ctaagcccac gctctctgag 2580
177 aaggccttgt atacactgaa ctccctcatg tatttctactg acaactgttt tctgtgcccc 2640
179 tgaagagttg caagagcacc ttgaaaacac aagcaaacat atcatatcaa cttccaggca 2700
181 tctaattaat tgtgcaccaa aatcgttacc cttccaatct tgcaacagaa gagcaaaagc 2760
183 tgcttagacc ttactcaaag gctccaaagc agaaacaagt tttgcttctt gcagcaatgt 2820
185 gtaacagagg aaacagatgt cctggaagat cccatggtct gtttcagggtg agatggaacc 2880
187 aatgctgcac acgaagattc atctcaaatg aaatgcacag gaaaagccaa ttactttatg 2940
189 tgaataaaag aataaatccc taaagcagtg gttctcaacc agagtgatac caccacacc 3000
191 ccagagggca tttgggaatt cttggggaca ttttggggtg acacactgaa ctgctggatg 3060
193 ctatcagcat ttagtaggta tgctcgatgt cttgcagaag gacatgatgg tcctacacag 3120
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201 ccaactacaa gtcataact gtagcatgga acctggtagt agcataataa ataaatTTTT 3360
203 agtaagaggc ttaagaaatt ttagcaaaaa aagcactccc tttcttctct cctacatctc 3420
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207 aagttcaaaa ccagatgctc attacagttc ttttaaacac caaactagtc atctcaaaaa 3540
209 tatggctaac tctctggact aaattccata ggaaaaatta ttaatttcaa aatgcctaata 3600
211 tttgatcaaa tgctgaaag agccaaaggc aatcatgtcc tgcttctcac tcagggcaga 3660
213 gttccatgag gtcagaaaag ctccaatgat atccggagggt ctgtcagaga ttaaaatctc 3720
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219 aaacaaacc caggctattt gcaggggggg ggaaagagat accccaaaag tcaaccctat 3900
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227 caattttcta gatgcctgac ctacagaca gacctgtttt cctcctgcca ggctttttt 4140
229 gccctccgc taggagcctg ggaaatcccc ctccaacctt ttgctggcct gtcaccagaa 4200
231 gttcataaaa tgaattcatc cagattctct gaatctctct tttcccagc catgtgcca 4260
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239 <211> LENGTH: 22
240 <212> TYPE: DNA
241 <213> ORGANISM: Homo sapiens
243 <400> SEQUENCE: 3
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248 <211> LENGTH: 21
249 <212> TYPE: DNA
250 <213> ORGANISM: Homo sapiens
252 <400> SEQUENCE: 4
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257 <211> LENGTH: 20
258 <212> TYPE: DNA
259 <213> ORGANISM: Homo sapiens
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267 <212> TYPE: DNA
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275 <211> LENGTH: 21
276 <212> TYPE: DNA
277 <213> ORGANISM: Homo sapiens
279 <400> SEQUENCE: 7
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283 <210> SEQ ID NO: 8
284 <211> LENGTH: 19
285 <212> TYPE: DNA
286 <213> ORGANISM: Homo sapiens
288 <400> SEQUENCE: 8
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292 <210> SEQ ID NO: 9
293 <211> LENGTH: 27
294 <212> TYPE: DNA
295 <213> ORGANISM: Artificial
297 <220> FEATURE:
298 <223> OTHER INFORMATION: Vector primer
300 <400> SEQUENCE: 9
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305 <211> LENGTH: 27
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307 <213> ORGANISM: Homo sapiens
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314 <211> LENGTH: 27
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323 <211> LENGTH: 27
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332 <211> LENGTH: 24
333 <212> TYPE: PRT
334 <213> ORGANISM: Homo sapiens
336 <400> SEQUENCE: 13

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339 1 5 10 15
342 Leu His Thr Lys Ile His Leu Lys
343 20
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347 <211> LENGTH: 56
348 <212> TYPE: PRT
349 <213> ORGANISM: Homo sapiens
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357 Trp Ile Thr Tyr Asn Ile Asn Ser Ser Leu Pro Tyr Thr Leu Leu Thr
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361 Pro Phe Pro Lys Gly Leu Ile Cys Ser Asn Leu Pro Val Pro Thr Val
362 35 40 45
365 Gln Trp Leu Ser Leu Pro Ser Pro
366 50 55
369 <210> SEQ ID NO: 15
370 <211> LENGTH: 78
371 <212> TYPE: PRT
372 <213> ORGANISM: Homo sapiens
374 <400> SEQUENCE: 15
376 Met Glu Pro Gly Ser Ser Ile Ile Asn Lys Phe Leu Val Arg Gly Leu
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380 Arg Asn Phe Ser Lys Lys Ser Thr Pro Phe Leu Pro Pro Tyr Ile Ser
381 20 25 30
384 His Met Phe Phe Asn Thr Lys Asn Ser Val Ile Leu Glu Lys Leu Leu
385 35 40 45
388 Thr Val Leu Leu Ser Ser Lys Pro Asp Ala His Tyr Ser Ser Phe Lys
389 50 55 60
392 His Gln Thr Ser His Leu Lys Asn Met Ala Asn Ser Leu Asp
393 65 70 75
396 <210> SEQ ID NO: 16
397 <211> LENGTH: 30
398 <212> TYPE: PRT
399 <213> ORGANISM: Homo sapiens
401 <400> SEQUENCE: 16
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407 Pro Met Ile Ser Gly Gly Leu Ser Glu Ile Lys Ile Ser Ser
408 20 25 30
411 <210> SEQ ID NO: 17
412 <211> LENGTH: 21
413 <212> TYPE: PRT
414 <213> ORGANISM: Homo sapiens
416 <400> SEQUENCE: 17
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419 1 5 10 15
422 Gly Arg Pro Ser Thr

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RAW SEQUENCE LISTING ERROR SUMMARY
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Invalid <213> Response:

Use of "Artificial" only as "<213> Organism" response is incomplete,
per 1.823(b) of New Sequence Rules. Valid response is Artificial Sequence.

Seq#:9

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L:9 M:270 C: Current Application Number differs, Replaced Current Application No

L:9 M:271 C: Current Filing Date differs, Replaced Current Filing Date